

Figure 1.

Name: SW_ROD_KIME_MOUSE	oo	Len: 468	Check: 7988	Weight: 1.00
Name: SW_ROD_KIME_RAT	oo	Len: 468	Check: 7364	Weight: 1.00
Name: SW_HUM_KIME_HUMAN	oo	Len: 468	Check: 8275	Weight: 1.00
Name: SW_OTHER_KIME_PYRAB	oo	Len: 468	Check: 8911	Weight: 1.00
Name: SW_OTHER_KIME_PYRHO	oo	Len: 468	Check: 673	Weight: 1.00
Name: SW_OTHER_KIME_PYRFU	oo	Len: 468	Check: 8129	Weight: 1.00
Name: SW_OTHER_KIME_ARATH	oo	Len: 468	Check: 9149	Weight: 1.00
Name: SW_OTHER_KIME_METTH	oo	Len: 468	Check: 6345	Weight: 1.00
Name: SW_OTHER_KIME_ARCFU	oo	Len: 468	Check: 5101	Weight: 1.00
Name: SW_OTHER_KIME_AERPE	oo	Len: 468	Check: 3583	Weight: 1.00
Name: SW_OTHER_KIME_SCHPO	oo	Len: 468	Check: 9129	Weight: 1.00
Name: SW_OTHER_KIME_YEAST	oo	Len: 468	Check: 1853	Weight: 1.00
Name: SW_OTHER_KIME_METJA	oo	Len: 468	Check: 8449	Weight: 1.00
Name: PARACOCUS	oo	Len: 468	Check: 7087	Weight: 1.00

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SW_ROD_KIME_MOUSE	..MLSEALLV	SAPGKVLHG	EHAVVHGKVA	LAAALN.LRT	FLLLRP....
SW_ROD_KIME_RAT	..MLSEVLLV	SAPGKVLHG	EHAVVHGKVA	LAVALN.LRT	FLVLRP....
SW_HUM_KIME_HUMAN	..MLSEVLLV	SAPGKVLHG	EHAVVHGKVA	LAVSLN.LRT	FLRLQP....
SW_OTHER_KIME_PYRAB	..MPRLVLA	SAPAKIILFG	EHSVYVGKPA	IASAID.LRT	YVRAEF....
SW_OTHER_KIME_PYRHO	..MVKYVLA	SAPAKVILFG	EHSVYVGKPA	IASAIE.LRT	YVRAQF....
SW_OTHER_KIME_PYRFU	....MKVIA	SAPAKVILFG	EHSVYVGKPA	IAAID.LRT	FVEAEL....
SW_OTHER_KIME_ARATH	....MEVKA	RAPGKILAG	EHAVVHGSTA	VAAID.LYT	YVTLRFPLPS
SW_OTHER_KIME_METTH	....MKSSA	SAPAKIILFG	EHAVVYKPA	IAAID.RRV	TVTUSE....
SW_OTHER_KIME_ARCFU	....MIA	SAPGKILFG	EHAVVYGRHA	VVSAIN.LRC	RVSVRK....
SW_OTHER_KIME_AERPE	..MRRAARA	SAPGKVIIVG	EHFVVRGSLA	IVAAIG.RRL	RVTVRS....
SW_OTHER_KIME_SCHPO	..MSKSLIV	SSPGKILFG	EHAVVYGATA	LAAAVS.LRS	YCKLOT....
SW_OTHER_KIME_YEAST	..MSLPFLT	SAPGKVIIFG	EHSVYKPA	VAASVSALRT	YLLISE....
SW_OTHER_KIME_METJA	.....MII	ETPSKVLFG	EHAVVYGYRA	ISMAID.LTS	TIEKETQ...
PARACOCUS	MSTGRPEAGA	HAPGKLILSG	EHSVLYGAPA	LAMAIA.RYT	EVWFTP....

Numbering

SW_ROD_KIME_MOUSE	.QSNKGVSUN	LPNIGIKQVW	DVGML...QR	LDTSFLEQGD	VSVPTLE.QL
SW_ROD_KIME_RAT	.QSNKGVSUN	LPNIGIKQVW	DVATL...QL	LDTSFLEQGD	VPAPTLE.QL
SW_HUM_KIME_HUMAN	.HNSGKVDLS	LPNIGIKRAW	DVARL...QS	LDTSFLEQGD	VTTPTSE.QV
SW_OTHER_KIME_PYRAB	.NDSGNKIE	AHDIKTP...	.....G	LIVSFSSED..	.KIYFET.DY
SW_OTHER_KIME_PYRHO	.NDSGNKIE	AHDIKTP...	.....G	LIVSFSSED..	.KIYFET.DY
SW_OTHER_KIME_PYRFU	.IREKKIRIE	AHDIKVP...	.....G	LTVSFSSEN..	.EYFET.DY
SW_OTHER_KIME_ARATH	AENNDRLTLQ	LKDISLEPSW	SLARIKEAIP	YDSSTLCR..	.STPASC.SE
SW_OTHER_KIME_METTH	...SSSTHVT	IPSLGIR...	.....H	SSER.....	.....
SW_OTHER_KIME_ARCFU	...SDRFLI	RSSLGES...	.....G	LDYQ.....	.....
SW_OTHER_KIME_AERPE	..GGKGIVLE	SSMLGRHS..	.....AP	LPGQ.....	.....
SW_OTHER_KIME_SCHPO	.TNNNEIVIV	MSDIGTERRW	N.....LQS	LPWQHVTVEN	VQHPASSPNL
SW_OTHER_KIME_YEAST	SSAPDTIELD	FPDISFNHKW	SINDFNAITE	DQVNSQKLAK	AQOATDGLSQ
SW_OTHER_KIME_METJA	...EDEIILN	LNDLNKS...	.....LG	LNLNEIKN..	.INPN...NF
PARACOCUS	LGIGEGIRTT	FANLSGGATY	S.....LK	LLSGFKSRLD	RRFEQFLNGD

Numbering

SW_ROD_KIME_MOUSE	EKLKKMGDLP	RD.RAGNEGM	ALLA...FLY	LYLAICRKQR	TLPSLDMVW
SW_ROD_KIME_RAT	EKLKKVAGLP	RD.CVNEGL	SLLA...FLY	LYLAICRKQR	TLPSLDMVW
SW_HUM_KIME_HUMAN	EKLKEVAGLP	DD.CAVTERL	AVLA...FLY	LYLSICRKQR	ALPSLDIVW
SW_OTHER_KIME_PYRAB	GKAAEVLSYV	R.....	..HA...IEL	VLEEADKR..	..TGVSVSIT
SW_OTHER_KIME_PYRHO	GKAAEVLSYV	R.....	..YA...IEL	ALEESDKR..	..VGIDVSIT
SW_OTHER_KIME_PYRFU	GKAAEVLSYV	R.....	..EA...INL	VLEEADKKN..	..VGIKVSIT
SW_OTHER_KIME_ARATH	ETLKSIAVLV	EEQNLPKEKM	WLSS...GIS	TFLWLYTRII	GENPATVVIN
SW_OTHER_KIME_METTH	P.SGGILDYI	G.....	...R...CLE	LYHDAS....	...PLDIRVE
SW_OTHER_KIME_ARCFU	R.HPXVQAV	K.....	.....	RFGELRN...	.IPGAEIEIE
SW_OTHER_KIME_AERPE	GAAAKVSPVL	EP.....	.....YIA	VLRSLAARGV	SVVPHTILVE
SW_OTHER_KIME_SCHPO	DLLOGLGELL	KNEENGLIHS	AMLC...TLY	LFTSLSSPS..	..QGCTLTIS
SW_OTHER_KIME_YEAST	ELVSLDPLL	AQLSESFHYH	AAFC...FLY	MFVCLCPHA..	..KNIKFSLK
SW_OTHER_KIME_METJA	GDFKYCLCAI	KN.....	.....TL	DYLNIEPK..	..TGFKINIS
PARACOCUS	LKVHKVLTHP	DDLAVYALAS	LLHDKPPGTA	AMPGIGAMHH	LPRPGELGSR

Numbering

Figure 1 (continued)

SW_ROD_KIME_MOUSE	SELPPGAGLG	SSAAYSVCVA	AALLTACEEV	SNPLKDGVS	SRWPEEDLKS
SW_ROD_KIME_RAT	SELPPGAGLG	SSAAYSVCVA	AALLTACEEV	SNPLKDRGSI	GSWPEEDLKS
SW_HUM_KIME_HUMAN	SELPPGAGLG	SSAAYSVCVA	AALLTVCEEI	PNPLKDGDCV	NRWTKEDLEL
SW_OTHER_KIME_PYRAB	SQIPVGAGLG	SSAAVAVATI	GAUSKLLDLE	LS.....	...KEE....
SW_OTHER_KIME_PYRHO	SQIPVGAGLG	SSAAVAVATI	GAUSKLLDLE	LS.....	...KEE....
SW_OTHER_KIME_PYRFU	SQIPVGAGLG	SSAAVAVATI	GAUSKLLDLE	LS.....	...KEE....
SW_OTHER_KIME_ARATH	SELPYGSGLG	SSAALCVALT	AALLASSISE	KTR...GNGW	SSLDETNLLEL
SW_OTHER_KIME_METTH	MEIPAGSGLG	SSAALTVALI	GALDRYHGRD	HG.....	...PGE....
SW_OTHER_KIME_ARCFU	SEIPIGSGLG	SSAAVIVATI	AALNAEFDGD	MD.....	...KEA....
SW_OTHER_KIME_AERPE	SGIPPRAGLG	SSAASMVAYA	LSYSAMHGDP	LS.....	...AED....
SW_OTHER_KIME_SCHPO	SQVPLGAGLG	SSATISVVVA	TSLLAFGNI	EPP...SSN.	SLQNNKALAL
SW_OTHER_KIME_YEAST	STLPIGAGLG	SSASISVSLA	LAMAYLGLI	GS.....NDL	EKLSENDKHI
SW_OTHER_KIME_METJA	SKIPISCGLG	SSASITIGTI	KAVSGFYNKE	LK.....	...DDE....
PARACOCUS	TELPIGAGMG	SSAAIVAATT	VLFTLLDRP	KT.....	...PEQ....

Numbering

SW\_ROD\_KIME\_MOUSE  
SW\_ROD\_KIME\_RAT  
SW\_HUM\_KIME\_HUMAN  
SW\_OTHER\_KIME\_PYRAB  
SW\_OTHER\_KIME\_PYRHO  
SW\_OTHER\_KIME\_PYRFU  
SW\_OTHER\_KIME\_ARATH  
SW\_OTHER\_KIME\_METTH  
SW\_OTHER\_KIME\_ARCFU  
SW\_OTHER\_KIME\_AERPE  
SW\_OTHER\_KIME\_SCHPO  
SW\_OTHER\_KIME\_YEAST  
SW\_OTHER\_KIME\_METJA  
PARACOCUS  
Numbering

INKWAFEGEER VIHGNPSGVD NAVSTWGGAL RFQ....QGT ..MSSLKSLP  
INKWAYEGER VIHGNPSGVD NSVSTWGGAL RYQ....QGK ..MSSLKRLP  
INKWAFQGER MIHGNPSGVD NAVSTWGGAL RYH....QGK ..ISSLKRSF  
IAKMGHKVEL LVQGASSGID PTVSAIGGFL VYK....QGE ..FEHLP.FV  
IAKLGHKVEL LVQGASSGID PTVSAVGGFL VYK....QGK ..FEHLP.FM  
IAKMGHKTTEL LVQGASSGID PTVSAIGGFI FYE....KGK ..FEHLP.FM  
LNKWAPEGEK IINGKPSGID NTVSAYGNMI KFC....SGE ..ITRLQSNM  
TAARAHREVEV DVQGAASPLD TAISTYGGVL YLDS....QRR ..VRQFE.AD  
IFQMAKQVEI DVQGRASGID PFISTFGGSW LFP....ERR ..KVEMP...  
LYSVAMEGEK IAHGKPSGVD VTIARVGGVL AYR....RGE NPVDIRPGLT  
IEAWSFLGEC CIHGTPSGID NAVATNGGLI AFR....KAT AHQSAMKEFL  
VNQWAFIGEK CIHGTPSGID NAVATYGNAL LFEKDSHNGT INTNNFKFLD  
IAKLGVMVEK EIQQKASITD TSTITYKGIL EIKNN...KFR KIKGEFEEFL  
RFDRVRFCER LKHGKAGPID AASVVRGGLV RVGGNG.PGS ISSFDLPEDH  
174 222

SW\_ROD\_KIME\_MOUSE  
SW\_ROD\_KIME\_RAT  
SW\_HUM\_KIME\_HUMAN  
SW\_OTHER\_KIME\_PYRAB  
SW\_OTHER\_KIME\_PYRHO  
SW\_OTHER\_KIME\_PYRFU  
SW\_OTHER\_KIME\_ARATH  
SW\_OTHER\_KIME\_METTH  
SW\_OTHER\_KIME\_ARCFU  
SW\_OTHER\_KIME\_AERPE  
SW\_OTHER\_KIME\_SCHPO  
SW\_OTHER\_KIME\_YEAST  
SW\_OTHER\_KIME\_METJA  
PARACOCUS  
Numbering

....SLQILL TNTKV.PRST KALVAVRSR L.TKFPEIVA PLLTSIDAIS  
....ALQILL TNTKV.PRST KALVAGVRSR L.IKFPEIMA PLLTSIDAIS  
....ALQILL TNTKV.PRNT RALVAGVRNR L.LKFPEIVA PLLTSIDAIS  
....ELPIVV GYTGS.SGST KELVAMVRRR Y.EEMPELIE PILESMGKLV  
....ELPIVV GYTGS.TGST KELVAMVRKR Y.EEMPELVE PILEAMGKLV  
....ELPIVV GYTGS.SGPT KELVAMVRKR Y.EEMPELIV PILEAMGKVV  
....PLRMLI TNTRV.GRNT KALVSGVSQR A.VRHPDAMK SVFNAVDSIS  
....LGDLYI AHLDY.SGET ARMVAGVAER F.RRFPDIMG RIMDTVESIT  
....FKFFV INFG..SRST AEMVAKVAEL R.ERHPEVVD KIFDAIDAIS  
....GVTLIV ADTGV.ERRT RDVVEHVLIS A.DALGEAST YIYRAADLIA  
KPKDTLSVMI TDTKQ.PKST KKLVOGVFEL K.ERLPTVID SIIDAIDGIS  
DFP.AIPMIL TYTRI.PRST KDLVARVRVL VTEKFPEVMK PILDAMGECA  
K...NCKFLI VYAEKRKKKT AELVNEVAKI E.....NKD EIFKEIDKVI  
DLVAGRGWYW VLHGRVSGT GECVSAVAAA H...G..RDA ALWDAPAVCT  
223 267

SW\_ROD\_KIME\_MOUSE  
SW\_ROD\_KIME\_RAT  
SW\_HUM\_KIME\_HUMAN  
SW\_OTHER\_KIME\_PYRAB  
SW\_OTHER\_KIME\_PYRHO  
SW\_OTHER\_KIME\_PYRFU  
SW\_OTHER\_KIME\_ARATH  
SW\_OTHER\_KIME\_METTH  
SW\_OTHER\_KIME\_ARCFU  
SW\_OTHER\_KIME\_AERPE  
SW\_OTHER\_KIME\_SCHPO  
SW\_OTHER\_KIME\_YEAST  
SW\_OTHER\_KIME\_METJA  
PARACOCUS  
Numbering

LECERVLGEM VAAP..... ..VPEQYLV LEELIDMNQH HLNALGVGHN  
LECERVLGEM AAAP..... ..VPEQYLV LEELMDMNQH HLNALGVGHA  
LECERVLGEM GEAP..... ..APEQYLV LEELIDMNQH HLNALGVGHA  
DKAKEVIISK LDE..... ..EEKFLK LGELMNINHG LLDALGVSTK  
DKAKEIILSK LDE..... ..EEKLTK LGELMNINHG LLDALGVSTK  
EKAKDVILSN VDK..... ..EEKFER LGVLMNINHG LLDALGVSTK  
KELAAIIQSK DETS..... ..VTEKEER IKELMEMNQ LLSMGVSHS  
NTAYRELLRN NTEP..... ..LGELMNINQH LLDALGVSTR  
LEASDVG..S AER..... ..LEELIATNQS LLRAIGVSNP  
REALHAIE.K GDA..... ..ER LGLIMNAAQG LLSSLGASSL  
KSAVLALTSE SDK..... ..NSSAKK LGEFIVLNQK LLECLGVSHY  
LQGLEIMTKL SKCKGTDDEA VETNNELYEQ LLELIRINHG LLVSIGVSHP  
DEALKIK..N KED..... ..FGKLMTKNHE LLKKNISTP  
RALEAALLSG GSP..... ..DAAITENQR LLERIGVSPA  
268 299

Figure 1 (continued)

SW\_ROD\_KIME\_MOUSE  
SW\_ROD\_KIME\_RAT  
SW\_HUM\_KIME\_HUMAN  
SW\_OTHER\_KIME\_PYRAB  
SW\_OTHER\_KIME\_PYRHO  
SW\_OTHER\_KIME\_PYRFU  
SW\_OTHER\_KIME\_ARATH  
SW\_OTHER\_KIME\_METTH  
SW\_OTHER\_KIME\_ARCFU  
SW\_OTHER\_KIME\_AERPE  
SW\_OTHER\_KIME\_SCHPO  
SW\_OTHER\_KIME\_YEAST  
SW\_OTHER\_KIME\_METJA  
PARACOCUS  
Numbering

SLDQLCQVTA AHG.LHSKLT GAG.....GG GCGITLLKPG LEQATVEAAK  
SLDQLCQVTA AHG.LHSKLT GAG.....GG GCGITLLKPG LERAKVEAAK  
SLDQLCQVTR ARG.LHSKLT GAG.....GG GCGITLLKPG LEQPEVEATK  
KLSELVYAAR TAGAIGAKLT GAG.....GG GC.MYALAPG KQRE....VA  
KLSELVYAAR TAGAIGAKLT GAG.....GG GC.MYALAPG RQRE....VA  
KLSELVYAAR VAGALGAKIT GAG.....GG GC.MYALAPN KQRE....VA  
SIEAVILTTV KHK.LVSKLT GAG.....GG GCVLTLTPTG TVVDK...VV  
ELSMVMYEAR NAGAAGSKIT GAG.....GG GS.IIAHCPG CVDD....VV  
EIDRTIAELE RMG.LNAKIT GAG.....GG GC.IFGLFKG EKPK.....LE  
EIETLVYRMR SAGALGAKLT GAG.....WG GCVIGLFKEG EVERG...LE  
SIDRVLQATK SIG..WTKLT GAG.....GG GCTITLLTPE CKEEEFKLCK  
GLELIKNSLD DLRIGSTKLT GAG.....GG GCSLTLLRRD ITQEIDSFK  
KLDRIVDIGN RFG.FGAKLT GAG.....GG GCVIILVNEE KEKE.....  
ATQALVAQIE EAG.GAAKIC GAGSVRGDGH GAVLVRIIDA QAMASVMARH  
300 348

SW\_ROD\_KIME\_MOUSE  
SW\_ROD\_KIME\_RAT  
SW\_HUM\_KIME\_HUMAN  
SW\_OTHER\_KIME\_PYRAB  
SW\_OTHER\_KIME\_PYRHO  
SW\_OTHER\_KIME\_PYRFU  
SW\_OTHER\_KIME\_ARATH  
SW\_OTHER\_KIME\_METTH  
SW\_OTHER\_KIME\_ARCFU  
SW\_OTHER\_KIME\_AERPE  
SW\_OTHER\_KIME\_SCHPO  
SW\_OTHER\_KIME\_YEAST  
SW\_OTHER\_KIME\_METJA  
PARACOCUS  
Numbering

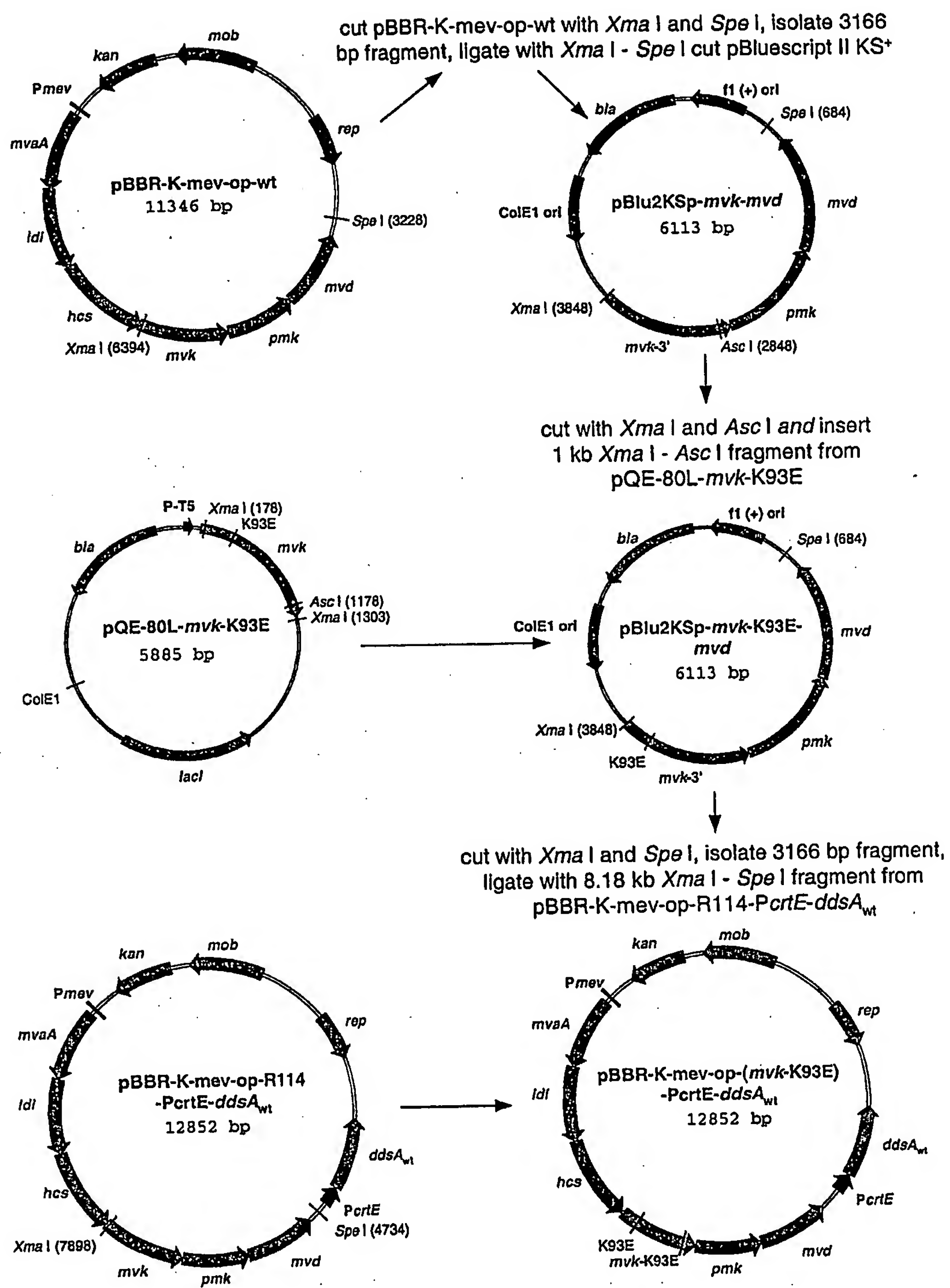
QALTSCG.FD CWETSIGAPG VSTHSAAVAVG DPVRQAL.GL .....  
QALTGCG.FD CWETSIGAPG VSMHSATSIE DPVRQAL.GL .....  
QALTSCG.FD CLETSIGAPG VSIHSATSLD SRVQQALDGL .....  
TAIKIAG.GT PMITRISKEG LRIEEVRE.. .....  
TAIKIAG.GI PMITRVSREG LRIEEVSR.. .....  
TAIRIAG.GT PMITEISREG LKIEEVIK.. .....  
EELESSG.FQ CFTALIGGNG AQICY..... .....  
TALNRN..WK AMRAEFSVKG LI..... .....  
.....G SFIVEPEKEG VRIEE..... .....  
SVVESS..Q AFTASIAEEG ARLEEF..... .....  
ESLLAHK.NS IYDVQLGGPG VSVVTDSDSF FPQYESDFDF KKLNLKSKFN  
KKLQDDFSYE TFETDLGGTG CCLLSAKNLN KDLKIKSLVF QLFENKTTK  
.....LLKELNKED VRIFNCRMMN .....  
PDLWAPLRM SRTGAAPGPA PRAQPLPGQG .....  
349 378

SW\_ROD\_KIME\_MOUSE  
SW\_ROD\_KIME\_RAT

.....  
.....

SW_HUM_KIME_HUMAN	.....
SW_OTHER_KIME_PYRAB	.....
SW_OTHER_KIME_PYRHO	.....
SW_OTHER_KIME_PYRFU	.....
SW_OTHER_KIME_ARATH	.....
SW_OTHER_KIME_METTH	.....
SW_OTHER_KIME_ARCFU	.....
SW_OTHER_KIME_AERPE	.....
SW_OTHER_KIME_SCHPO	KYYI.....
SW_OTHER_KIME_YEAST	QQIDDL LLPG NTNLPWTS
SW_OTHER_KIME_METJA	.....
PARACOCUS	.....

Figure 2.



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